## SEQUENCE LISTING

<110> Lu et al. <120> Polypeptide <130> 69856 <150> PCT/GB03/04296 <151> 2003-10-03 <150> 0306261.9 <151> 2003-03-19 <150> 0223193.4 <151> 2002-10-07 <160> 9 <170> PatentIn version 3.1 <210> 1 <211> 6 <212> PRT <213> Artificial sequence <220> <223> p53 inhibitor peptide <400> 1 Gly Pro Glu Glu Thr Asp <210> 2 <211> 7 <212> PRT <213> artificial peptide <220> <223> p53 inhibitor peptide <400> 2 Asp Gly Pro Glu Glu Thr Asp 1 5 <210> 3 <211> 6 <212> PRT <213> artificial peptide <220> <223> p53 inhibitor peptide

<400> 3 Thr Thr Leu Ser Asp Gly <210> <211> 6 <212> PRT <213> artificial peptide <220> <223> p53 inhibitor peptide <400> 4 Pro Arg Asn Tyr Phe Gly <210> 5 <211> 18 <212> PRT <213> artificial peptide <220> <223> p53 inhibitor peptide <400> 5 Arg Leu Gln Pro Ala Leu Pro Pro Glu Ala Gln Ser Val Pro Glu Leu 10 15 Glu Glu <210> 6 <211> 2156 <212> DNA <213> Homo sapiens <400> 6 geggeeget egaceeggeg tteagaegeg ggeagetace ggegeteget gggeteegeg 60 gggccgtcgg gcactttgcc tcgcagctgg cagcccgtca gccgcatccc catgcccccc 120 tecageeece ageeeegegg ggeeeegege cagegteeca teceeeteag catgatette 180 aagetgeaga aegeettetg ggageaeggg geeageegeg ceatgeteee tgggteeeee 240 ctcttcaccc gagcaccccc gcctaagctg cagccccaac cacaaccaca gccccagcca 300 caatcacaac cacageeeca getgeeecaa cageeecaga cecaaeeeca aaceeetaee 360 ccagcetece acateegeat ecceaacaga catggeecee tgtgaacgaa ggaeceecea 420

aacccccac cgagctggag cctgagccgg agatagaggg gctgctgaca ccagtgctgg 480 540 aggctggcga tgtggatgaa ggaccctgta gcaaggcctc tcagccccac gaggctgcag 600 ccagcactgc caccggaggc acagtcggtg cccgagctgg aggaggtggc acgggtgttg geggaaatte eeeggeeeet caaacgeagg ggeteeatgg ageaggeeee tgetgtggee 660 ctgccccta cccacaagaa acagtaccag cagatcatca gccgcctctt ccatcgtcat 720 780 ggggggccag ggcccggggg gcggagccag agctgtcccc catcactgag ggatctgagg ccagggcagg gcccctgct cctgcccac cagctccat tccaccgccc ggccccgtcc 840 cagagcagcc caccagagca gccgcagagc atggagatgc gctctgtgct gcggaaggcg 900 ggctccccgc gcaaggcccg ccgcgcgcc ctcaaccctc tggtgctcct cctggacgcg 960 1020 gcgctgaccg gggagctgga ggtggtgcag caggcggtga aggagatgaa cgacccgagc 1080 cagcccaacg aggagggcat cactgccttg cacaacgcca tctgcggcgc caactactct atcgtggatt tcctcatcac cgcgggtgcc aatgtcaact cccccgacag ccacggctgg 1140 acaccettge actgegegge gtegtgeaac gacacagtea tetgeatgge getggtgeag 1200 . caeggegetg caatettege caecaegete agegaeggeg ceaecgeett egagaagtge 1260 gaccettace gegagggtta tgetgactge gecacetace tggcagacgt egagcagagt 1320 atggggctga tgaacagcgg ggcagtgtac gctctctggg actacagcgc cgagttcggg 1380 gacgagetgt cetteegega gggegagteg gteacegtge tgeggaggga egggeeggag 1440 1500 gagaccgact ggtggtgggc cgcgctgcac ggccaggagg gctacgtgcc gcggaactac 1560 ttcgggctgt tccccagggt gaagcctcaa aggagtaaag tctagcagga tagaaggagg 1620 tttctgaggc tgacagaaac aagcattcct gccttccctc cagacctctc cctctgtttt ttgctgcctt tatctgcacc cctcaccctg ctggtggtgg tccttgccac cggttctctg 1680 ttctcctgga agtccaggga agaaggaggg ccccagcctt aaatttagta atctgcctta 1740 gccttgggag gtctgggaag ggctggaaat cactggggac aggaaaccac ttccttttgc 1800 caaatcagat cccgtccaaa gtgcctccca tgcctaccac catcatcaca tcccccagca 1860 1920 agccagccac ctgcccagcc gggcctggga tgggccacca caccactgga tattcctggg agtcactgct gacaccatct ctcccagcag tcttggggtc tgggtgggaa acattggtct 1980 ctaccaggat ccctgcccca cctctcccca attaagtgcc ttcacacagc actggtttaa 2040 tgtttataaa caaaatagag aaactggttt aatgtttata aacaaaatag agaaactttc 2100

gcttataaat aaaagtagtt	tgcacagaaa	tgaaaaaaaa	aaaaaaaaa	aaaaaa	2156
<210> 7 <211> 2310 <212> DNA <213> Caenorhabditis	elegans				
<400> 7				<b>h</b> - <b>h</b>	60
atggtcacga ccagtagcgg					60
gtgtctctga ttcactcgtc	ggattctgta	cgaactgttt	caactgcccc	aatataccgt	120
ccgacgtcat caatggcatc	tacgatggct	cataaatctt	cgacggctcc	gttcatctcc	180
gcaaatcaac gaatgtcaaa	accgccggtt	cgggtggtcg	ctcaaccacc	accaccacat	240
ccacaagcat tgtcccaaca	gtatcaccag	cagaatccga	tgatgatgta	ttccgcacca	300
aatacacgac cacacgttat	tccgacaatg	caagtgcaac	cgacaatggc	cgctcaaatt	360
aaacgaaata atcctgttaa	tgcacagttt	cagaaccctt	ctgaaatgat	cgccgattac	420
ggtgtaaaac cgcagtcagt	agaaatggtg	caaagagttc	gagctgttcg	aagacaagtc	480
gccgacgagg agaccgaact	gcgaagactc	agagagcttg	aacacgaaac	ggcacagctt	540
caaaataaga attatggaag	agaaagagag	ttgaatgtgc	aaggatccat	gctgaaagaa	600
gctcaattag agttgagaaa	tgcttcaatg	agggcgcaat	ctttaaacaa	gcatttggaa	660
gaaatgtacc ggagaagaca	aactgcagca	gcggcagcgc	tcgtggaaca	acgaaaaatg	720
cagcaacatc agattcttct	agcccgagct	gcaaatcaag	tatccacaca	agaagttata	780
agacctcgtg cttctgtcga	accattccaa	gttaataata	cccaacagca	acaaccatca	840
cctcaaatga tgaaatcaga	agaattttcg	gagaaaagag	atttgaatgg	acaaactggc	900
agttatgatg ctatcgatgg	atcaggagat	catcaaaaaa	taccgacgga	gccatcgtac	960
ttggcaccat gtaaagaaaa	ccagcaaaaa	tactcggagt	taagtaaaat	ggcatctacg	1020
gatcctcatt caaatcacag	ttcaccatca	acttcttcgc	agaaagctcc	gacgttgatc	1080
acattttctc caccaagttt	tgaacagaaa	atcaactcgt	ctacaatgac	tcgggattct	1140
ccgttcgttg agcgtccaac	atcgtttggt	gatagtctag	acgaatcacg	actgagaagt	1200
ggaaagactg atttggtatc	acttcgatca	gattccctga	aagctacgaa	acgtcgttct	1260
tgggctgctt ccgaaggtac	ttcaatgtca	gaggcagaga	tgattcatag	gcttcttgat	1320
gaacaacgtc gtgggagatc	acattttatt	ccacaattgc	caacatcaca	agaagaacca	1380
tcggcaataa catcagaaac	atatgccgaa	gaagttgtca	attcagaatc	gaaacaagtt	1440

gctacaagtt cggattccac	taataatctt	gaattgccaa	ccgaacaaat	ggtattaggt	1500
agtgatacca caacagaaga	agatgcaagt	tcgtgttcaa	cacgttctga	tgatggacag	1560
aatcttgaaa tggaagttgc	gattgaaaga	agaactgtta	aaggcatttt	gagaagacct	1620
aatgaaaaga tgaacaaagg	tcgcattgaa	tttgacccat	tagcactctt	gctcgatgct	1680
gctttagaag gagaactcga	tttagtgaga	agcagtgcct	caaagctaac	agatgtctca	1740
caggccaatg atgaagggat	tacggcgttg	cacaatgcga	tttgtgctgg	acactatgag	1800
attgtaagat ttttgatcga	gaacgacgct	gatgtgaatg	ctcaagattc	cgatggttgg	1860
actccacttc attgtgcagc	ttcctgtaat	aaccttccaa	tggttagaca	acttgtggaa	1920
ggaggaggat gcgttctcgc	ttcgacacta	tctgatatgg	aaacacctgt	ggagaagtgt	1980
gaagaagatg aagatggtta	tgatggatgt	ttgaagtatc	tttccgcagc	ccataactca	2040
acgggatcaa ttaatactgg	aaaagtttac	gctgcttatg	gatatgaggc	ggcatttgaa	2100
gatgagctca gttttgatgc	aggagatgaa	ttgacggtta	ttgagaaaga	taaagtcgat	2160
aaaaattggt ggacatgtga	gaagaacaat	ggagagaagg	gacaagtacc	aagaacatat	2220
ttggcgttgt acccatcgtt	aaaatacaga	aagaagctca	actttgtgat	gttcgatctt	2280
ccattggaat cgaacaacaa	tgtcgaataa				2310

<210> 8

<211> 350

<212> PRT

<213> Homo sapiens

<400> 8

Met Trp Met Lys Asp Pro Val Ala Arg Pro Leu Ser Pro Thr Arg Leu 1 5 10 15

Gln Pro Ala Leu Pro Pro Glu Ala Gln Ser Val Pro Glu Leu Glu Glu 20 25 30

Val Ala Arg Val Leu Ala Glu Ile Pro Arg Pro Leu Lys Arg Arg Gly 35 40 45

Ser Met Glu Gln Ala Pro Ala Val Ala Leu Pro Pro Thr His Lys Lys 50 60

Gln Tyr Gln Gln Ile Ile Ser Arg Leu Phe His Arg His Gly Gly Pro 65 70 75 80

Arg Pro Gly Gln Gly Pro Leu Leu Pro His Gln Leu Pro Phe His Arg Pro Ala Pro Ser Gln Ser Ser Pro Pro Glu Gln Pro Gln Ser Met Glu Met Arg Ser Val Leu Arg Lys Ala Gly Ser Pro Arg Lys Ala Arg Arg Ala Arg Leu Asn Pro Leu Val Leu Leu Asp Ala Ala Leu Thr Gly Glu Leu Glu Val Val Gln Gln Ala Val Lys Glu Met Asn Asp Pro Ser Gln Pro Asn Glu Glu Gly Ile Thr Ala Leu His Asn Ala Ile Cys Gly Ala Asn Tyr Ser Ile Val Asp Phe Leu Ile Thr Ala Gly Ala Asn Val Asn Ser Pro Asp Ser His Gly Trp Thr Pro Leu His Cys Ala Ala Ser Cys Asn Asp Thr Val Ile Cys Met Ala Leu Val Gln His Gly Ala Ala Ile Phe Ala Thr Thr Leu Ser Asp Gly Ala Thr Ala Phe Glu Lys Cys Asp Pro Tyr Arg Glu Gly Tyr Ala Asp Cys Ala Thr Tyr Leu Ala Asp Val Glu Gln Ser Met Gly Leu Met Asn Ser Gly Ala Val Tyr Ala Leu Trp Asp Tyr Ser Ala Glu Phe Gly Asp Glu Leu Ser Phe Arg Glu

Gly Pro Gly Gly Arg Ser Gln Ser Cys Pro Pro Ser Leu Arg Asp Leu

Gly Glu Ser Val Thr Val Leu Arg Arg Asp Gly Pro Glu Glu Thr Asp 305 310 315 320

Trp Trp Trp Ala Ala Leu His Gly Gln Glu Gly Tyr Val Pro Arg Asn 325 330 335

Tyr Phe Gly Leu Phe Pro Arg Val Lys Pro Gln Arg Ser Lys 340 345 350

<210> 9

<211> 769

<212> PRT

<213> Caenorhabditis elegans

<400> 9

Met Val Thr Thr Ser Ser Gly Gly Gly Ile Gly Tyr Pro Ala Asn Asn 1 5 10 15

Gly Val Thr Gln Val Ser Leu Ile His Ser Ser Asp Ser Val Arg Thr
20 25 30

Val Ser Thr Ala Pro Ile Tyr Arg Pro Thr Ser Ser Met Ala Ser Thr 35 40 45

Met Ala His Lys Ser Ser Thr Ala Pro Phe Ile Ser Ala Asn Gln Arg 50 55 60

Met Ser Lys Pro Pro Val Arg Val Val Ala Gln Pro Pro Pro Pro His 65 70 75 80

Pro Gln Ala Leu Ser Gln Gln Tyr His Gln Gln Asn Pro Met Met Met 85 90 95

Tyr Ser Ala Pro Asn Thr Arg Pro His Val Ile Pro Thr Met Gln Val
100 105 110

Gln Pro Thr Met Ala Ala Gln Ile Lys Arg Asn Asn Pro Val Asn Ala 115 120 125

Gln Phe Gln Asn Pro Ser Glu Met Ile Ala Asp Tyr Gly Val Lys Pro 130 135 140

Gln Ser Val Glu Met Val Gln Arg Val Arg Ala Val Arg Arg Gln Val 145 150 155 160

Ala Asp Glu	Glu Thr 165	Glu Leu	Arg .	_	Leu 170	Arg	Glu	Leu	Glu	His 175	Glu
Thr Ala Gln	Leu Gln 180	Asn Lys		Tyr ( 185	Gly	Arg	Glu	Arg	Glu 190	Leu	Asn
Val Gln Gly 195	Ser Met	Leu Lys	Glu . 200	Ala (	Gln	Leu	Glu	Leu 205	Arg	Asn	Ala
Ser Met Arg 210	Ala Gln	Ser Leu 215	Asn	Lys l	His	Leu	Glu 220	Glu	Met	Tyr	Arg
Arg Arg Gln 225	Thr Ala	Ala Ala 230	Ala	Ala 1		Val 235	Glu	Gln	Arg	Lys	Met 240
Gln Gln His	Gln Ile 245	Leu Leu	Ala	_	Ala 250	Ala	Asn	Gln	Val	Ser 255	Thr
Gln Glu Val	Ile Arg 260	Pro Arg		Ser V 265	Val	Glu	Pro	Phe	Gln 270	Val	Asn
Asn Thr Gln 275	Gln Gln	Gln Pro	Ser 280	Pro (	Gln	Met	Met	Lys 285	Ser	Glu	Glu
Phe Ser Glu 290	Lys Arg	Asp Leu 295	Asn	Gly (	Gln	Thr	Gly 300	Ser	Tyr	Asp	Ala
Ile Asp Gly 305	Ser Gly	Asp His 310	Gln :	Lys :		Pro 315	Thr	Glu	Pro	Ser	Tyr 320
Leu Ala Pro	Cys Lys 325	Glu Asn	Gln		Lys 330	Tyr	Ser	Glu	Leu	Ser 335	Lys
Met Ala Ser	Thr Asp 340	Pro His		Asn I 345	His	Ser	Ser	Pro	Ser 350	Thr	Ser
Ser Gln Lys 355	Ala Pro	Thr Leu	Ile '	Thr I	Phe	Ser	Pro	Pro 365	Ser	Phe	Glu
Gln Lys Ile 370	Asn Ser	Ser Thr 375	Met '	Thr <i>l</i>	Arg	Asp	Ser 380	Pro	Phe	Val	Glu

Arg Pro Thr Ser Phe Gly Asp Ser Leu Asp Glu Ser Arg Leu Arg Ser Gly Lys Thr Asp Leu Val Ser Leu Arg Ser Asp Ser Leu Lys Ala Thr Lys Arg Arg Ser Trp Ala Ala Ser Glu Gly Thr Ser Met Ser Glu Ala Glu Met Ile His Arg Leu Leu Asp Glu Gln Arg Arg Gly Arg Ser His Phe Ile Pro Gln Leu Pro Thr Ser Gln Glu Glu Pro Ser Ala Ile Thr Ser Glu Thr Tyr Ala Glu Glu Val Val Asn Ser Glu Ser Lys Gln Val Ala Thr Ser Ser Asp Ser Thr Asn Asn Leu Glu Leu Pro Thr Glu Gln Met Val Leu Gly Ser Asp Thr Thr Thr Glu Glu Asp Ala Ser Ser Cys Ser Thr Arg Ser Asp Asp Gly Gln Asn Leu Glu Met Glu Val Ala Ile Glu Arg Arg Thr Val Lys Gly Ile Leu Arg Arg Pro Asn Glu Lys Met Asn Lys Gly Arg Ile Glu Phe Asp Pro Leu Ala Leu Leu Leu Asp Ala Ala Leu Glu Gly Glu Leu Asp Leu Val Arg Ser Ser Ala Ser Lys Leu Thr Asp Val Ser Gln Ala Asn Asp Glu Gly Ile Thr Ala Leu His Asn Ala Ile Cys Ala Gly His Tyr Glu Ile Val Arg Phe Leu Ile Glu Asn 

Asp Ala Asp Val Asn Ala Gln Asp Ser Asp Gly Trp Thr Pro Leu His 610 615 620

Cys Ala Ala Ser Cys Asn Asn Leu Pro Met Val Arg Gln Leu Val Glu 625 630 635 640

Gly Gly Cys Val Leu Ala Ser Thr Leu Ser Asp Met Glu Thr Pro 645 650 655

Val Glu Lys Cys Glu Glu Asp Glu Asp Gly Tyr Asp Gly Cys Leu Lys 660 665 670

Tyr Leu Ser Ala Ala His Asn Ser Thr Gly Ser Ile Asn Thr Gly Lys 675 680 685

Val Tyr Ala Ala Tyr Gly Tyr Glu Ala Ala Phe Glu Asp Glu Leu Ser 690 695 700

Phe Asp Ala Gly Asp Glu Leu Thr Val Ile Glu Lys Asp Lys Val Asp 705 710 715 720

Lys Asn Trp Trp Thr Cys Glu Lys Asn Asn Gly Glu Lys Gly Gln Val
725 730 735

Pro Arg Thr Tyr Leu Ala Leu Tyr Pro Ser Leu Lys Tyr Arg Lys Lys 740 745 750

Leu Asn Phe Val Met Phe Asp Leu Pro Leu Glu Ser Asn Asn Asn Val 755 760 765

Glu